

SEQUENCE LISTING

3H CENTER 1600/2900

<110> McBurney, Robert N.
Holt, William
Gwynne, David I.
Marchionni, Mark

<120> Therapeutic Methods Comprising Use of a Neuregulin

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<140> US 09/530,884

<141> 2000-08-29

<150> PCT/US98/21349

<151> 1998-10-08

<150> US 60/062,109

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Val Val Ile Glu Gly Lys Val His Pro Gln Arg Arg Gln Gln Gly Ala
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gcc Ala 225	tct Ser	ttc Phe	ccc Pro	cct Pro	ctg Leu 230	gag Glu	acg Thr	ggc Gly	cgg Arg	aac Asn 235	ctc Leu	aag Lys	aag Lys	gag Glu	gtc Val 240	720
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Gly	Asp	Arg 115		Pro	Pro	Ala	Ala 120		Pro	Arg	Ala	Leu 125	Gly	Pro	Pro	
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Cys	GIÀ	Arg	35	пÃр	GIU	Азр	261	40	ıyı	116	1110	1110	45	OI u	110	
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Ser Asp Ser Ser Gly Ser Arg Met Gly Met Asn * Ile Glu Lys Thr
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Tyr Met Cys Lys Val Ile Ser Lys Leu Gly Asn Asp Ser Ala Ser Ala
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Asn Ile Thr Ile Val Glu Ser Asn Ala
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ctg Leu	gct (	gat Asp	tct Ser	gga Gly 15	gaa Glu	tat Tyr	atg Met	tgc Cys	aaa Lys 20	gtg Val	atc Ile	agc Ser	aaa Lys	cta Leu 25	gga Gly	161
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	tgt Cys 60				tgaa	atcac	ege a	aggto	gtgtg	ja aa	atcto	attg:	, tga	aacaa	ata	312
aaaa tcac	tcat tcca	ga a ta a	agga aggto	aaad gaaat	ct ct ta ga	atgt accto	ttga gaaat	a aat c ata	atct	tat agat	gggt tatt	cctc	ct q	gtaaa	agctct	372 417
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tca gag tct ccc a Ser Glu Ser Pro I	tt aga ata tca le Arg Ile Ser 20	gta tcc aca gaa Val Ser Thr Glu 25	gga gca aat act Gly Ala Asn Thr 30	95
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His Pro Asn Pro Pro Pro Glu Asn Val Gln Leu Val Asn Gln Tyr Val
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ctt Leu	cgg Arg 50	tct Ser	gaa Glu	aga Arg	aac Asn	acc Thr 55	atg Met	atg Met	aac Asn	gta Val	gcc Ala 60	aac Asn	Gly ggg	ccc Pro	cac His	192
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act Thr	gtc Val	act Thr 115	cag Gln	act Thr	ccc Pro	agt Ser	cac His 120	agc Ser	tgg Trp	agc Ser	aat Asn	gga Gly 125	cac His	act Thr	gaa Glu	384
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cat His 65	cct Pro	aac Asn	cca Pro	ccc Pro	ccc Pro 70	gag Glu	aat Asn	gtc Val	cag Gln	ctg Leu 75	gtg Val	aat Asn	caa Gln	tac Tyr	gta Val 80	240
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				gaa Glu												432
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Val Glu Glu Arg Pro Leu Leu Leu Val Thr Pro Pro Arg Leu Arg
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Glu Lys Tyr Asp His His Ala Gln Gln Phe Asn Ser Phe His Cys Asn
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Pro Ala His Glu Ser Asn Ser Leu Pro Pro Ser Pro Leu Arg Ile Val
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Glu Asp Glu Glu Tyr Glu Thr Thr Gln Glu Tyr Glu Pro Ala Gln Glu
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Pro Val Lys Lys Leu Thr Asn Ser Ser Arg Arg Ala Lys Arg Thr Lys
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Pro Asn Gly His Ile Ala His Arg Leu Glu Met Asp Asn Asn Thr Gly
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Ala Asp Ser Ser Asn Ser Glu Ser Glu Thr Glu Asp Glu Arg Val Gly
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Glu Asp Thr Pro Phe Leu Ala Ile Gln Asn Pro Leu Ala Ala Ser Leu
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                                                       175
Glu Ala Ala Pro Ala Phe Arg Leu Val Asp Ser Arg Thr Asn Pro Thr
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His Thr Pro Ser Ser Pro Lys Ser Pro Pro Ser Glu Met Ser Pro Pro
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gag Glu 65	aag Lys	tat Tyr	gac Asp	cac His	cac His 70	gcc Ala	cag Gln	caa Gln	ttc Phe	aac Asn 75	tcg Ser	ttc Phe	cac His	tgc Cys	aac Asn 80	241
ccc Pro	gcg Ala	cat His	gag Glu	agc Ser 85	aac Asn	agc Ser	ctg Leu	ccc Pro	ccc Pro 90	agc Ser	ccc Pro	ttg Leu	agg Arg	ata Ile 95	gtg Val	289
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	gct Ala 210	aac Asn	caa Gln	gac Asp	cct Pro	atc Ile 215	gct Ala	gtc Val	taaa	accg	aa a	taca.	.ccca	ıt		672
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Ser Gln Ser Pro
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135

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tac Tyr	tcc Ser	tct Ser 115	ctc Leu	aag Lys	ttc Phe	aag Lys	tgg Trp 120	ttc Phe	aag Lys	aat Asn	GJA aaa	agt Ser 125	gaa Glu	tta Leu	agc Ser	384
cga Arg	aag Lys 130	aac Asn	aaa Lys	cca Pro	gaa Glu	aac Asn 135	atc Ile	aag Lys	ata Ile	cag Gln	aaa Lys 140	agg Arg	ccg Pro	ggg ggg	aag Lys	432

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at Me	g tgo t Cys	c aaa E Lys	a gtg 5 Val	atc Ile 165	agc Ser	aaa Lys	cta Leu	gga Gly	aat Asn 170	gac Asp	agt Ser	gcc Ala	tct Ser	gcc Ala 175	aac Asn	528
ato Ile	e acc	att Ile	gtg Val 180	gag Glu	tca Ser	aac Asn	gcc Ala	aca Thr 185	tcc Ser	aca Thr	tct Ser	aca Thr	gct Ala 190	Gly	aca Thr	576
ago Sei	cat His	Leu 195	gtc Val	aag Lys	tgt Cys	gca Ala	gag Glu 200	aag Lys	gag Glu	aaa Lys	act Thr	ttc Phe 205	tgt Cys	gtg Val	aat Asn	624
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Gly	Arg	Leu 35	Lys	Glu .	Asp :				Ile	Phe		Met (	30 Glu	Pro (	Glu	
	50		Ser			Pro (	Gly A				Ser 60	Leu :				
65			Gly		70					Gly 75	Gln				R O	
Gln	Arg	Cys	Ala :	Leu	Pro 1	Pro 1	Arg 1	Leu :	Lys (	Glu	Met	Lys :	Ser (	Gln (	Glu	

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tca Ser	tcc Ser	Tnr	tcc Ser 170	aca Thr	tct Ser	aca Thr	gct Ala	ggg Gly 175	aca Thr	agc Ser	cat His	Leu	gtc Val 180	aag Lys	tgt Cys	1004
gca Ala	GIU	aag Lys 185	gag ( Glu )	aaa Lys	act Thr	ttc Phe	tgt Cys 190	gtg Val	aat Asn	gga Gly	ggc Gly	gag Glu 195	tgc Cys	ttc Phe	atg Met	1052
var.	aaa Lys 200	gac Asp	ctt : Leu :	tca Ser	Asn	ccc Pro 205	tca Ser	aga Arg	tac Tyr	Leu	tgc Cys 210	aag Lys (	tgc Cys	cca Pro .	aat Asn	1100
gag Glu 215	ttt a Phe '	act ( Thr (	ggt q Gly 1	Asp A	cgc Arg 220	tgc Cys	caa Gln .	aac Asn	Tyr	gta Val 225	atg ( Met /	gcc a Ala :	agc Ser	Phe '	tac Tyr 230	1148
agt a Ser 1	acg f	tcc a Ser '	rnr i	ecc Pro 1 235	ttt Phe	ctg Leu	tct ( Ser 1	Leu	cct Pro 240	gaa Glu	tagg	cgcai	tg c	tcag	tcggt	1201
actto tttci atgao gacao	cetet cgaaa caata gteed gtaca	tg to at to at ct at ac	egact ecgtg gatet ggeet etett ettga	geen jacta tgaa tgaa tata	a gtg a tta a aag a aag	gcct; gggc actg gtct; atga; ggta;	gteg tetg tgat caet ceet agtt	age acg ttta atce	gaga tact acat attg	aca cgt gat aga	ttaad aggto agtod aaata	cacaa gegta cetet aaaaa	ag contag grant ag	gatte gctce cccae gttce	ttacc gtatg cagtg gtgca cacgg ttgta ctgac	1321 1381 1441 1501

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